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(54) Title: DETECTION OF EPIGENETIC ABNORMALITIES AND DIAGNOSTIC METHOD BASED THEREON

Sch74-E52m/Sch74-E51m CAGCTCACTGCRAACCTCCGCC-TCCCTGGATTCAAGC-GATTTTCCGCCCTTAG-CCTCTTGACTAACTGGACTAGAGGCCAGGTACCACAGGCCAGCTAATTTT-GTATT
BD43-E76m/BD43-E83m CGGCTCATGCAACCTCGGCC-TCCCTGGTTCAAGC-AATTCTCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
BD34-D19N TGGCTCATGTYAACCTCTGCC-TCCCTGGTTCAAGC-AATTCTCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
BD34-E62m-B BD43-15m TGGCTCATGTYAACCTCTGCC-TCCCTGGTTCAAGC-AATTCTCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
Ctr157-E5m CGCTCATGCAACCTCGGCC-TCCCTGGTTCAAGC-GATTCCTCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
BD43-RevE7m CAGCTCATGCAACCCACACCG-TCCCTGGTTCAAGC-GATTATCCCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
BD43-Rev7m CAGCTCATGCAACCCACACCG-TCCCTGGTTCAAGC-GATTATCCCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
BD34-A14M CAGCCCCAGCQAACCTCGGCC-TCCCTGGTTCAAGC-GATTATCCCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
Ctr157-E3m CGGCTCATGCAACCTCGGCC-TCCCTGGTTCAAGC-GATTATCCCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
Sch74-E318m Sch74-E318m TGGCTCATGCAACCTCTGCC-TCCCTGGTTCAAGC-AATTCTCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
Sch74-E318_m TGGCTCATGCAACCTCTGCC-TCCCTGGTTCAAGC-AATTCTCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
Ctr157-Z4m CAGCTCATGCAACCTCTGCC-TCCCTGGTTCAAGC-GATTCCTCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
BD43-E79m CAGCTCATGCAACCTCTGCC-TCCCTGGTTCAAGC-GATTCCTCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
Ctr157-E6m CAGCTCATGCAACCTCTGCC-TCCCTGGTTCAAGC-GATTCCTCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
Ctr150-RevE169m CAGCTCATGCAACCTCTGCC-TCCCTGGTTCAAGC-GATTCCTCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
Sch56-E23m CAGCTCATGCAACCTCTGCC-TCCCTGGTTCAAGC-GATTCCTCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
Sch56-r-37m CAGCTCATGCAACCTCTGCC-TCCCTGGTTCAAGC-GATTCCTCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
Sch56-E23m CAGCTCATGCAACCTCTGCC-TCCCTGGTTCAAGC-GATTCCTCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
Ctr150-E16m CAGCTCATGCAACCTCTGCC-TCCCTGGTTCAAGC-GATTCCTCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT
Ctr150-E249m CAGCTCATGCAACCTCTGCC-TCCCTGGTTCAAGC-GATTCCTCTCGTCAG-CCTCCCGAGTAGCTGGGATCACGGCACATGCCAACTAATTTT-GTATT

- (57) Abstract: The present invention provides a method of detecting an epigenetic abnormality associated with a disease. The method comprises identifying, within a eukaryotic genome, a locus having a hypomethylated sequence specific for the disease and an endogenous multi-copy DNA element. The method can also comprise separate steps of identifying a disease-specific hypomethylated sequence and identifying an endogenous multi-copy DNA element, where the steps may be performed in any order, so long as a locus is identified that has both a disease-specific hypomethylated sequence and an endogenous multi-copy DNA element. The disease-specific hypomethylated sequences detected in accordance with the present invention indicate putative regions of epigenetic dys-regulation and indicate aberrantly regulated nucleic acid sequences that may cause or predispose a patient to disease, such as, but not limited to, Huntington's disease, cancers, diabetes, schizophrenia, or bipolar disorder.

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